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(54) Title: ENDODERM, CARDIAC AND NEURAL INDUCING FACTORS

#### (57) Abstract

Novel proteins have been designated "cerberus" and "frzb-1", respectively. Cerebus is expressed as a secreted peptide during embryogenesis of the Xenopus embryo, and is expressed specifically in the head organizer region. This new molecule has endodermal, cardiac, and neural tissue inducing activity, that should prove useful in therapeutic, diagnostic, and clinical applications requiring regeneration, differentiation, or repair of these and other tissues. Frzb-1 is a soluble antagonist of growth factors of the Wnt family that acts by binding to Wnt growth factors in the extracellular space. A third novel protein is termed PAPC which promotes the formation of dorsal mesoderm and somites in the embryo.

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# ENDODERM, CARDIAC AND NEURAL INDUCING FACTORS

#### 5 Field of the Invention

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The invention generally relates to growth factors, neurotrophic factors, and their inhibitors, and more particularly to several new growth factors with neural, endodermal, and cardiac tissue inducing activity, to complexes and compositions including the factors, and to DNA or RNA coding sequences for the factors. Further, one of the novel growth factors should be useful in tumor suppression gene therapy.

This application claims the benefit of U.S.

Provisional Application No. 60/020,150, filed June 20,
1996.

This invention was made with Government support under grant contract number HD-21502, awarded by the National Institutes of Health. The Government has certain rights in this invention.

#### Background of the Invention

Growth factors are substances, such as polypeptide hormones, which affect the growth of defined populations of animal cells in vivo or in vitro, but which are not nutrient substances. Proteins involved in the growth and differentiation of tissues may promote or inhibit growth, and promote or inhibit differentiation, and thus the general term "growth factor" includes cytokines, trophic factors, and their inhibitors.

widespread neuronal cell death accompanies normal development of the central and peripheral nervous systems. Studies of peripheral target tissues during development have shown that neuronal cell death results from the competition among neurons for limiting amounts of survivor factors ("neurotrophic factors"). The earliest identified of these, nerve growth factor ("NGF"), is the most fully characterized and has been shown to be essential for the survival of sympathetic and neural crest-derived sensory neurons during early development of both chick and rat.

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One family of neurotropic factors are the Wnts, which have dorsal axis-inducing activity. Most of the Wnt proteins are bound to cell surfaces. (See, e.g., Sokol et al., Science, 249, pp. 561-564, 1990.) Dorsal axis-inducing activity in Xenopus embryos by one member of this family (Xwnt-8) was described by Smith and Harland in 1991, Cell, 67, pp. 753-765. The authors described using RNA injections as a strategy for identifying endogenous RNAs involved in dorsal patterning to rescue dorsal development in embryos that were ventralized by UV irradiation.

Another member of the growth and neurotropic factor family was subsequently discovered and described by Harland and Smith, which they termed "noggin." (Cell, 70, pp. 829-840 (1992).) Noggin is a good candidate to function as a signaling molecule in by virtue of Nieuwkoop's center, its transcripts, and in Spemann's organizer, through its zygotic organizer-specific expression. Besides noggin, other secreted factors may be involved in the organizer phenomenon.

Another Xenopus gene designated "chordin" that begins to be expressed in Spemann's organizer and that can completely rescue axial development in ventralized embryos was described by Sasai et al., Cell, 79, pp. 779-790, 1994. In addition to dorsalizing mesoderm, chordin has the ability to induce neural tissue and its activities are antagonized by Bone Morphogenetic Protein-4 (Sasai et al., Nature, 376, pp. 333-336, 1995).

Therefore, the dorsal lip or Spemann's organizer of the Xenopus embryo is an ideal tissue for seeking novel growth and neurotrophic factors. New growth and neurotrophic factors are useful agents, particularly those that are secreted due to their ability to be used in physiologically active, soluble forms because these factors, their receptors, and DNA or RNA coding sequences therefore and fragments thereof are useful in a number of therapeutic, clinical, research, diagnostic, and drug design applications.

#### Summary of the Invention

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In one aspect of the present invention, the sequence the novel peptide that can substantially purified form is shown by SEQ ID NO:1. The Xenopus derived SEQ ID NO:1 has been designated -"cerberus," and this peptide is capable of inducing endodermal, cardiac, and neural tissue development in vertebrates when expressed. The nucleotide sequence which. when expressed results in cerberus. illustrated by SEQ ID NO:2. Since peptides of the invention induce endodermal, cardiac, and neural tissue differentiation in vertebrates, they should be able to be prepared in physiologically active form for a number of therapeutic, clinical, and diagnostic applications.

Cerberus was isolated during a search for molecules expressed specifically in Spemann's organizer containing a secretory signal sequence. In addition to cerberus, two other novel cDNAs were identified.

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Xenopus derived peptide that can deduced from SEQ ID NO:3 encodes a novel protein we had earlier designated as "frazzled," a secreted protein of 318 amino acids that has dorsalizing activity in Xenopus We now designate the novel protein as embryos. "frzb-1." The gene for frzb-1 is expressed in many adult tissues of many animals, three of the cDNAs (Xenopus, mouse, and human) have been cloned by us. accession numbers for the Xenopus, mouse, and human frzb-1 cDNA sequences of the gene now designated frzb-1 are U68059, U68058, and U68057, respectively. has some degree of sequence similarity to the Drosophila gene frizzled which has been shown to encode a seventransmembrane protein that can act both as a signalling and as a receptor protein (Vinson et al., Nature, 338, pp. 263-264, 1989; Vinson and Adler, Nature, 329, pp. 549-551, 1987). Vertebrate homologues of Frizzled have been isolated and they too were found to be anchored to the cell membrane by seven membrane spanning domains (Wang et al., J. Biol. Chem., 271, pp. 4468-4476, 1996). Frzb-1 differs from the frizzled proteins in that it is an entirely soluble, diffusible secreted protein and therefore -suitable - as a -therapeutic -agent. The nucleotide sequence derived from Xenopus that, when expressed, results in frzb-1 protein is illustrated by SEO ID NO:4. The frzb-1 protein derived from mouse is shown as SEQ ID-NO:7, while the mouse frzb-1 nucleotide sequence is SEQ ID NO:8. The human derived frzb-1 protein is illustrated by SEQ ID NO:9, and the human frzb-1 nucleotide sequence is SEQ ID NO:10.

Frzb-1 is an antagonist of Whts in vivo, and thus is believed to find utility as a tumor suppressor gene, since overexpressed Wht proteins cause cancer. Frzb-1 may also be a useful vehicle for solubilization

and therapeutic delivery of Wnt proteins complexed with it.

The final cDNA isolated containing a signal sequence results in a peptide designated Paraxial Protogadherin (PAPC). The cDNA for PAPC is a divergent member of the cadherin multigene family. PAPC is most related to protocadherin 43 reported by Sano et al., The EMBO J., 12, pp. 2249-2256, 1993. As shown in SEQ ID NO:5, the PAPC gene encodes a transmembrane protein of 896 amino acids, of which 187 are part intracellular domain. PAPC is a cell adhesion molecule, and microinjection of PAPC mRNA constructs into Xenopus embryos suggest that PAPC acts as a molecule involved in mesoderm differentiation. A soluble form of the PAPC extracellular domain is able to block muscle and mesoderm formation in Xenopus embryos. The nucleotide sequence encoding Xenopus PAPC is provided in SEO ID NO:6.

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Cerberus, frzb-1, or PAPC or fragments thereof (which also may be synthesized by in vitro methods) may be fused (by recombinant expression or in vitro covalent methods) to an immunogenic polypeptide and this, in turn, may be used to immunize an animal in order to raise antibodies against the novel proteins. Antibodies are recoverable from the serum of immunized animals. Alternatively, monoclonal antibodies may be prepared from-cells from the immunized animal in conventional fashion. Immobilized antibodies are useful particularly in the diagnosis (in vitro or in vivo) or purification of cerberus, frzb-1, or PAPC.

Substitutional, deletional, or insertional mutants of the novel polypeptides may be prepared by in vitro or recombinant methods and screened for immunocrossreactivity with cerberus, frzb-1, or PAPC and for cerberus antagonist or agonist activity.

Cerberus or frzb-1 also may be derivatized in vitro in order to prepare immobilized and labelled proteins, particularly for purposes of diagnosis of insufficiencies thereof, or for affinity purification of antibodies thereto.

Among applications for the novel proteins are tissue replacement therapy and, because frzb-1 is an antagonist of Wnt signaling, tumor suppression therapies. The cerberus receptor may define a novel signalling pathway. In addition, frzb-1 could permit the isolation of novel members of the Wnt family of growth factors.

#### Brief Description of the Drawings

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Figure 1 illustrates the amino acid sequence (SEQ ID NO:1) of the Fig. 2 cDNA clone for cerberus;

Figure 2 illustrates a cDNA clone (SEQ ID NO:2) for cerberus derived from Xenopus. Sense strand is on top (5' to 3' direction) and the antisense strand on the bottom line (in the opposite direction);

Figures 3 and 4 show the amino acid and nucleotide sequence, respectively, of full-length frzb-1 from Xenopus (SEQ ID NOS:3 and-4);

Figures 5 and 6 show the amino acid and nucleotide sequence, respectively, of full-length PAPC from Xenopus (SEQ ID NOS:5 and 6);

Figures 7 and 8 show the amino acid and nucleotide sequence, respectively, of full-length frzb-1 from mouse (SEQ ID NOS:7 and 8); and

Figures 9 and 10 show the amino acid and nucleotide sequence, respectively, of full-length frzb-1 from human (SEQ ID NOS:9 and 10).

#### Detailed Description of the Preferred Embodiments

Among the several novel proteins and their nucleotide sequences described herein, is a novel endodermal, cardiac, and neural inducing factor in vertebrates that we have named "cerberus." When \_\_referring\_to cerberus,\_the present invention also contemplates the use of fragments, derivatives, agonists, or antagonists of cerberus molecules. cerberus has no homology to any reported growth factors, 10 it is proposed to be the founding member of a novel family of growth factors with potent biological activities, which may be isolated using SEQ ID NO:2.

The amphibian organizer consists of several cell populations with region-specific activities. On the basis of morphogenetic movements, three very different cell populations can distinguished in the organizer. First, cells with crawling migration movements involute, fanning out to form the prechordal plate. Second, cells involute through the dorsal lip driven by convergence and extension movements, giving rise to the notochord of the trunk. Third, involution ceases and the continuation of -mediolateral-intercalation movements leads to posterior extension movements and to the formation of the tail notochord and of the chordoneural hinge. The three cell populations correspond to the head, trunk, and tail -organizers, respectively. ----

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The cerberus gene is expressed at the right time and place to participate in cell signalling by Spemann's organizer. Specifically, cerberus is expressed in the head organizing region that consists of crawling-migrating cells. The cerberus expressing region corresponds to the prospective foregut, including the liver and pancreas anlage, and the heart mesoderm.

Cerberus expression is activated by chordin, noggin, and organizer-specific homeobox genes.

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Our studies were conducted in early embryos of the frog Xenopus laevis. The frog embryo is well suited to experiments, particularly experiments pertaining to generating-and-maintaining regional differences within the embryo for determining roles in tissue differentiation. It is easy to culture embryos with access to the embryos even at very early stages of development (preceding and during the formation of body pattern and 10 differentiation) and the embryos are large. The initial work with noggin and chordin also had been in Xenopus embryos, and, as predicted, was highly conserved among vertebrates. Predictions based on work with Xenopus as to corresponding human noggin were proven true and the ability to clone the gene for human noggin was readily accomplished. (See the description of Xenopus work and cloning information in PCT application, published March 17, 1994, WO 9 405 800, and the subsequent human cloning based thereon in the PCT application, also published March 17, 1994, as WO 9 405 791.)

#### CLONING

The cloning of cerberus, frzb-1, and PAPC resulted from a comprehensive screen for cDNAs enriched Spemann's organizer. Subtractive differential screening was performed as follows. In brief, poly A<sup>+</sup> RNA was isolated from 300 dorsal lip and ventral marginal zone (VMZ) explants at stage 10%. After first strand cDNA synthesis approximately 70-80% of common sequences were removed by substraction with biotinylated VMZ poly A+ RNA prepared from 1500 ventral gastrula For differential screening, duplicate filters (2000 plaques per 15 cm plate, a total of 80,000 clones

screened) of an unamplified oriented dorsal lip library were hybridized with radiolabeled dorsal lip or VMZ Putative organizer-specific clones were isolated, grouped by sequence analysis from the 5' end and wholemount in situ hybridization, and subsequently classified into known and new dorsal-specific genes. Rescreening of the library (100,000 independent phages) with a probe resulted in the isolation cerberus additional clones, 31 of which had similar size as the longest one of the 11 original clones indicating that they were presumably full-length cDNAs. The longest cDNAs for cerberus, frzb-1, and PAPC were completely sequenced.

To explore the molecular complexity of

Spemann's organizer we performed a comprehensive
differential screen for dorsal-specific cDNAs. The
method was designed to identify abundant cDNAs without
bias as to their function. As shown in Table 1, five
previously known cDNAs and five new ones were isolated,
of which three (expressed as cerberus, frzb-1, and PAPC,
respectively) had secretory signal sequences.

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#### TABLE\_1

	Previously Known Genes	Gene Product	No. of Isolates
	Chordin	novel secreted protein	70
·	Goosecoid	homeobox gene	3
5	Pintallavis/XFKH-1	forkhead/transcription factor	2
	Xnot-2	homeobox gene	1
	Xlim-1	homeobox gene	1
•	New Genes		
	Cerberus	novel secreted protein	11
10	PAPC	cadherin-like/transmembrane	2
	Frzb-1	novel secreted protein	. 1
	Sox-2	sry/transcription factor	1
	Fkh-like	forkhead/transcription factor	1

The most abundant dorsal-specific cDNA was chordin (chd), with 70 independent isolates. The second most abundant cDNA was isolated 11 times and named cerberus (after a mythological guardian dog with multiple heads). The cerberus cDNA encodes a putative secreted polypeptide of 270 amino acids, with an amino terminal hydrophobic signal sequence and a carboxy terminal cysteine-rich region (Fig. 1). Cerberus is expressed specifically in the head organizer region of the Xenopus embryo, including the future foregut.

An abundant mRNA found in the dorsal region of the Xenopus gastrula encodes the novel putative secreted protein we have designated as cerberus. Cerberus mRNA has potent inducing activity in Xenopus embryos, leading to the formation of ectopic heads. Unlike other organizer-specific factors, cerberus does not dorsalize mesoderm and is instead an inhibitor of trunk-tail mesoderm. Cerberus is expressed in the anterior-most

domain of the gastrula including the leading edge of the deep layer of the dorsal lip a region that, as shown here, gives rise to foregut and midgut endoderm. Cerberus promotes the formation of cement gland, olfactory placodes, cyclopic eyes, forebrain, and duplicated heart and liver (a foregut derivative). Because the pancreas is also derived from this foregut region, it is likely that cerberus induces pancreas in addition to liver. The expression pattern and inducing activities of cerberus suggest a role for a previously neglected region of the embryo, the prospective foregut endoderm, in the induction of the anterior head region of the embryo.

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putative secreted protein transiently expressed during embryogenesis and the deduced amino acid sequence of Xenopus cerberus is shown. The signal peptide sequence and the nine cysteine residues in the carboxy-terminus are indicated in bold. Potential N-linked glycosylation sites are underlined. In database searches the cerberus protein showed limited similarity only to the mammalian Dan protein, a possible tumor suppressor proposed to be a DNA-binding protein.

its amino acid sequence and the spacing of its 9 cysteine residues were not significantly similar to other proteins in the databases (NCBI-Gen Bank release 93.0). We conclude that the second most abundant dorsal-specific cDNA encodes a novel putative secreted factor, which should be the founding member of a novel family of growth factors active in cell differentiation.

<u>Cerberus Demarcates an Anterior Organizer</u> <u>Domain</u>. Cerberus mRNA is expressed at low levels in the unfertilized egg, and zygotic transcripts start accumulating at early gastrula. Expression continues during gastrula and early neurula, rapidly declining during neurulation. Importantly, cerberus expression starts about one hour after that of chd, suggesting that cerberus could act downstream of the chd signal.

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Whole-mount in situ hybridizations reveal that expression starts in the yolky endomesodermal cells located in the deep layer of the organizer. The cerberus domain includes the leading edge of the most anterior organizer cells and extends into the lateral mesoderm. The leading edge gives rise to liver, pancreas, and foregut in its midline, and the more lateral region gives rise to heart mesoderm at later stages of development.

Fig. 2 sets out the sequence of a full length 15 Xenopus cDNA for cerberus.

This entirely new molecule has demonstrated physiological properties that should prove useful in therapeutic, diagnostic, and clinical applications that require regeneration, differentiation, or repair of tissues, such wound repair, neuronal regenerational or transplantation, supplementation of heart muscle differentiation, differentiation of pancreas and liver, and other applications in which cell differentiation processes are to be induced.

The second, novel, secreted protein we have discovered is called "frzb-1," which was shown to be a secreted protein in Xenopus oocyte microinjection experiments. Thus it provides a natural soluble form of the related extracellular domains of Drosophila and vertebrate frizzled proteins. We propose that the latter proteins could be converted into active soluble forms by introducing a stop codon before the first transmembrane domain. We have noted that the cysteinerich region of frzb-1 and frizzled contains some overall structural homology with Wnt proteins using the Profile

Search homology program (Gribskov, Meth. Enzymol., 183, pp. 146-159, 1990). This had raised the interesting possibility that frzb-1 could interact directly with Wnt growth factors in the extracellular space. because we had found that when microinjected into embryos, frzb-1 constructs have dorsalizing activity, leading to the formation of embryos with enlarged brain and head, and shortened Somatic muscle differentiation, which requires truck. Xwnt-8, was inhibited. In the case of frzb-1, an attractive hypothesis, suggested by the structural homologies, was that it may act as an inhibitor of Wnt-8, a growth factor that has ventralizing activity in the Xenopus embryo (Christian and Moon, Genes Dev., 7, pp. 13-28, 1993). We have shown that frzb-1 can interact with Xwnt-8 and Wnt-1, and it is expected that it could also interact with other members of the Wnt family of growth factors, of which at least 15 members exist in mammals. In addition, a possible interaction with Wnts was suggested by the recent discovery that dishevelled, a gene acting downstream of wingless, has strong genetic interaction with frizzled mutants in Drosophila (Krasnow et al., Development, 121, pp. 4095-4102, 1995). This possibility has been explored in depth (Leyns et al., Cell, 88, pp. 747-756, March 21, 1997), because a soluble antagonist of the Wnt family of -proteins is expected to be of great therapeutic value. Examples 1 and 2 illustrate tests that show antagonism of Xwnt-8 by binding to frzb-1.

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Vertebrate homologues of Frizzled have been isolated and they too are anchored to the cell membrane by seven membrane spanning domains (Wang et al., J. Biol. Chem., 271, pp. 4468-4476, 1996). Frzb-1 differs from the frizzled proteins in that it is an entirely soluble, diffusible secreted protein and

therefore suitable as a therapeutic agent. The nucleotide sequence that when expressed results in frzb-1 protein is illustrated by SEQ ID NO:4.

ID NO:4 corresponds to the SEO homolog, but by using it in BLAST searches (and by -- cloning mouse frzb-1) we had been able to assemble the sequence of the entire mature human frzb-1 protein, SEO Indeed, human frzb-1 is encoded in six expressed sequence tags (ESTs) available in Genebank. 10 The human frzb-1 sequence can be assembled overlapping in the 5' to 3' direction the ESTs with the following accession numbers in Genebank: R63748, W38677, W44760, H38379, and N71244. No function had yet been assigned to these EST sequences, but we believe and thus propose here that human frzb-1 will 15 have similar functions in cell differentiation to those described above for Xenopus frzb-1. The nucleotide sequence of human frzb-1 is shown in SEQ ID NO:10. mouse frzb-1 protein and nucleotide sequences are 20 provided by SEQ ID NOS:7 and 8, respectively.

In particular, we believe that frzb-1 will prove useful in gene therapy of human cancer cells. In this rapidly developing field, one approach is to introduce vectors expressing anti-sense sequences to block expression of dominant ocogenes and growth factor receptors. Another approach is to produce episomal vectors that will replicate in human cells in a controlled fashion without transforming the cells. For an example of the latter (an episomal expression vector system for human gene therapy), reference is made to U.S. Patent 5,624,820, issued April 29, 1997, inventor Cooper.

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Gene therapy now includes uses of human tumor suppression genes. For example, U.S. Patent 5,491,064, issued February 13, 1996, discloses a tumor suppression

gene localized on chromosome 11 and described as potentially useful for gene therapy in cancers deleted or altered in their expression of that gene. maps to chromosome 2q31-33 and loss of one copy of the 2q31-33 and loss of one copy of the 2q arm has been observed with high incidence in lung carcinomas, colo-rectal carcinomas, and neuroblastomas, which has lead to the proposal that the 2q arm carries a tumor suppressor gene. We expect frzb to be suppressor gene, and thus to be useful in tumor suppression applications.

A number of applications for cerberus and frzb-1 are suggested from their pharmacological (biological activity) properties.

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15 For example, the cerberus and frzb-1 cDNAs should be useful as a diagnostic tool (such as through use of antibodies in assays for proteins in cell lines or use of oligonucleotides as primers in a PCR test to amplify those with sequence similarities to the oligonucleotide primer, and to determine how much of the novel protein is present).

Cerberus, of course, might act upon its target cells via its own receptor.— Cerberus, therefore, provides the key to isolate this receptor. Since many receptors mutate to cellular oncogenes, the cerberus receptor should prove useful as a diagnostic probe for certain tumor types. Thus, when one views cerberus as ligand in complexes, then complexes in accordance with the invention include antibody bound to cerberus, antibody bound to peptides derived from cerberus, cerberus bound to its receptor, or peptides derived from cerberus bound to its receptor or other factors. Mutant forms of cerberus, which are either more potent agonists or antagonists, are believed to be clinically useful.

Such complexes of cerberus and its binding protein partners will find uses in a number of applications.

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Practice of this invention includes use of an oligonucleotide construct comprising a sequence coding for cerberus or frzb-1 and for a promoter sequence operatively linked in a mammalian or a viral expression Expression and cloning vectors contain a nucleotide sequence that enables the vector to replicate in one or more selected host cells. Generally, in cloning vectors this sequence is one that enables the replicate independently of vector to the chromosomes, and includes origins of replication or autonomously replicating sequences. The well-known plasmid pBR322 is suitable for most gram negative bacteria, the 2µ plasmid origin for yeast and various viral origins (SV40, polyoma, adenovirus, VSV or BPV) are useful for cloning vectors in mammalian cells.

Expression and cloning vectors should contain a selection gene, also termed a selectable marker. Typically, this is a gene that encodes a protein necessary for the survival or growth of a host cell transformed with the vector. The presence of this gene ensures that any host-cell which-deletes the vector will not obtain an advantage in growth or reproduction over transformed hosts. Typical selection genes encode proteins that (a) confer resistance to antibiotics or other toxins, e.g. ampicillin, neomycin, methotrexate or tetracycline, (b) complement auxotrophic deficiencies.

Examples of suitable selectable markers for mammalian cells are dihydrofolate reductase (DHFR) or thymidine kinase. Such markers enable the identification of cells which were competent to take up the cerberus nucleic acid. The mammalian cell transformants are placed under selection pressure which only the transformants are uniquely adapted to survive by virtue

of having taken up the marker. Selection pressure is imposed by culturing the transformants under conditions in which the concentration of selection agent in the Amplification is the medium is successively changed. process by which genes in greater demand for production of a protein critical for growth are tandem within reiterated in the chromosomes of . successive generations of recombinant cells. quantities of cerberus or frzb-1 can therefor be synthesized from the amplified DNA.

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For example, cells transformed with the DHFR selection gene are first identified by culturing all of the transformants in a culture medium which contains methotrexate (Mtx), a competitive antagonist of DHFR. An appropriate host cell in this case is the Chinese hamster ovary (CHO) cell line deficient activity, prepared and propagated as described by Urlaub and Chasin, Proc. Nat. Acac. Sci., 77, 4216 (1980). transformed cells then are exposed to increased levels This leads to the synthesis of multiple copies of the DHFR gene and, concomitantly, multiple copies of other DNA comprising the expression vectors, such as the -DNA- encoding-cerberus-or-frzb-1. --Alternatively, host cells transformed by an expression vector comprising DNA sequences encoding cerberus or frzb-1 and aminoglycoside 3' phosphotransferase (APH) protein can be selected by cell-growth in-medium-containing an aminoglycosidic antibiotic such as kanamycin or neomycin or G418. Because eukaryotic cells do not normally express an endogenous APH activity, genes encoding APH protein, commonly referred to as neo resistant genes, may be used as dominant selectable markers in a wide range of eukaryotic host cells, by which cells transformed by the vector can readily be identified.

Expression vectors, unlike cloning vectors, should contain a promoter which is recognized by the host organism and is operably linked to the cerberus nucleic acid. Promoters are untranslated sequences located upstream from the start codon of a structural -- gene -- (generally within about 100 to 1000 - bp) that control the transcription and translation of nucleic acid under their control. They typically fall into two constitutive. classes, inducible and promoters are promoters that initiate increased levels of transcription from DNA under their control response to some change in culture conditions, e.g. the presence or absence of a nutrient or a change in temperature. At this time a large number of promoters recognized by a variety of potential host cells are well These promoters can be operably linked to cerberus encoding DNA by removing them from their gene of origin by restriction enzyme digestion, followed by insertion 5' to the start codon for cerberus or frzb-1.

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Nucleic acid is operably linked when it is placed into a functional relationship with another DNA for nucleic acid sequence. For example, presequence or secretory leader is operably linked to DNA for a polypeptide if it is expressed as a preprotein which participates in the secretion of the polypeptide; a promoter or enhancer is operably linked to a coding sequence if it affects the transcription of the sequence; or a ribosome binding site is operably linked to a coding sequence if it is positioned so as to Generally, operably linked facilitate translation. means that the DNA sequences being linked are contiguous and, in the case of a secretory leader, contiguous and in reading phase. Linking is accomplished by ligation at convenient restriction sites. If such sites do not

exit then synthetic oligonucleotide adapters or linkers are used in accord with conventional practice.

Transcription of the protein-encoding DNA in mammalian host cells is controlled by promoters obtained from the genomes of viruses such as polyoma, cytomegalovirus, adenovirus, retroviruses, hepatitis-B virus, and most preferably Simian Virus 40 (SV40), or from heterologous mammalian promoters, e.g. the actin promoter. Of course, promoters from the host cell or related species also are useful herein.

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Cerberus and frzb-1 are clearly useful as a component of culture media for use in culturing cells, such as endodermal, cardiac, and nerve cells, in vitro. We believe cerberus and frzb-1 will find uses as agents for enhancing the survival or inducing the growth of liver, pancreas, heart, and nerve cells, such as in tissue replacement therapy.

The final cDNA isolated containing a signal sequence results in a peptide designated Paraxial Protocadherin (PAPC). The cDNA for PAPC is a divergent member of the cadherin multigene family. PAPC is most related to protocadherin 43 reported by Sano et al., The -EMBO J., 12, pp.-2249-2256, 1993. As-shown in SEQ ID NO:5, the PAPC gene encodes a transmembrane protein of amino acids, of which 187 are part intracellular domain. PAPC is a cell adhesion molecule, and-microinjection-of-PAPC-mRNA constructs-into-Xenopus embryos acts suggest that PAPC in mesoderm differentiation. The nucleotide sequence encoding Xenopus PAPC is provided in SEQ ID NO:6.

Therapeutic formulations of the novel proteins may be prepared for storage by mixing the polypeptides having the desired degree of purity with optional physiologically acceptable carriers, excipients or stabilizers, in the form of lyophilized cake or aqueous

solutions. Acceptable carriers, excipients stabilizers are nontoxic to recipients at the dosages and concentrations employed, and include buffers such as phosphate, citrate, and other organic acids; antioxidants including ascorbic acid; low molecular weight (less than about 10 residues) polypeptides; proteins, such as serum albumin, gelatin or immunoglobulins. components can include glycine, blutamine, asparagine, arginine, or lysine; monosaccharides, disaccharides, and other carbohydrates including glucose, mannose, or dextrins; chelating agents such as EDTA; sugar alcohols such as mannitol or sorbitol; saltforming counterions such as sodium; and/or nonionic surfactants such as Tween, Pluronics or PEG.

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Polyclonal antibodies to the novel proteins generally are raised in animals by multiple subcutaneous (sc) or intraperitoneal (ip) injections of cerberus or frzb-1 and an adjuvant. It may be useful to conjugate these proteins or a fragment containing the target amino acid sequence to a protein which is immunogenic in the immunized, be e.g., keyhole hemocyanin, serum albumin, bovine thyroglobulin, soybean - trypsin inhibitor using a bifunctional derivatizing agent, for example, maleimidobenzoyl sulfosuccinimide ester (conjugation through cysteine residues), N-hydroxysuccinimide (through residues), glutaraldehyde, succinic anhydride, SOCl2, or  $R^1N = C = NR$ .

Animals can be immunized against the immunogenic conjugates or derivatives by combining 1 mg or 1
µg of conjugate (for rabbits or mice, respectively)
with 3 volumes of Freund's complete adjuvant and
injecting the solution intradermally in multiple sites.
One month later the animals are boosted with 1/5 to 1/10
the original amount of conjugate in Fruend's complete

adjuvant by subcutaneous injection at multiple sites. Seven to 14 days later animals are bled and the serum is assayed for anti-cerberus titer. Animals are boosted until the titer plateaus. Preferably, the animal is boosted with the conjugate of the same cerberus or frzb-l\_polypeptide, but conjugated to a different protein and/or through a different cross-linking agent. Conjugates also can be made in recombinant cell culture as protein fusions. Also, aggregating agents such as alum are used to enhance the immune response.

Monoclonal antibodies are prepared by recovering spleen cells from immunized animals and immortalizing the cells in conventional fashion, e.g. by fusion with myeloma cells or by EB virus transformation and screening for clones expressing the desired antibody.

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Antibodies are useful in diagnostic assays for cerberus, frzb-1, or PAPC or their antibodies and to identify family members. In one embodiment of a receptor binding assay, an antibody composition which 20 binds to all of a selected plurality of members of the cerberus family is immobilized on an insoluble matrix, the test sample is contacted with the immobilized antibody composition in order to adsorb all cerberus 25 family members, and then the immobilized family members are contacted with a plurality of antibodies specific for—each member, each of the antibodies being individually identifiable as specific for a predetermined family member, as by unique labels such as 30 discrete fluorophores or the like. By determining the presence and/or amount of each unique label, the relative proportion and amount of each family member can be determined.

The antibodies also are useful for the 35 affinity purification of the novel proteins from

recombinant cell culture or natural sources. Antibodies that do not detectably cross-react with other growth factors can be used to purify the proteins free from these other family members.

#### EXAMPLE 1

Frzb-1 Antagonizes Xwnt-8 Non-Cell Autonomously

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To whether frzb-1 test can antagonize secondary axes caused by Xwnt-8 after secretion by injected cells, an experimental design was used. frzb-1 mRNA was injected into each of the four animal blastomeres of eight-cell embryos, and subsequently, a single injection of Xwnt-8 mRNA was given to a vegetalventral blastomere at the 16-32 cell stage. independent experiments, we found that injection of frzb-1 alone (n=13) caused mild dorsalization with 15 enlargement of the cement gland in all embryos and that injection of Xwnt-8 alone (n=53) lead to induction of complete secondary axes in 67% of the embryos. injection of frzb-1 into animal caps abolished the formation of complete axes induced by Xwnt-8 (n=27), 20 -leaving-only-a residual-14%-of-embryos-with very weak secondary axes. The double-injected embryos retained the enlarged cement gland phenotype caused by injection Because both mRNAs encode of frzb-1 mRNA alone. 25 —secreted proteins—and—were microinjected into different cells, we conclude that the antagonistic effects of frzb-1 and Xwnt-8 took place in the extracellular space after these proteins were secreted.

#### EXAMPLE 2

### Membrane-Anchored Wnt-1 Confers Frzb-1 Binding

To investigate a possible interaction between frzb-1 and Wnts, the first step was to insert an HA epitope tag into a Xenopus frzb-1 construct driven by the CMV (cytomegalovirus) promoter. Frzbl-HA was tested in mRNA microinjection assays in Xenopus embryos and found to be biologically active. Conditioned medium from transiently transfected cells contained up to  $10 \mu g/ml$  of Frzbl-HA (quantitated on Western blots using an HA-tagged protein standard).

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Transient transfection of 293 cells has been in demonstrating interactions between instrumental wingless and frizzled proteins. We therefore took advantage of constructs in which Wnt-1 was fused at the amino terminus of CD8, generating a transmembrane protein containing biologically active Wnt-1 exposed to the extracellular compartment. A Wnt1CD8 cDNA construct (a generous gift of Dr. H. Varmus, NIH) was subcloned into the pcDNA (Invitrogen) vector and transfected into 293 cells. After incubation with Frzbl-HA-conditioned -medium-(overnight at 37°C), -intensely labeled cells were observed by immunofluorescence. As a negative control, a construct containing 120 amino acids of Xenopus chordin, an unrelated secreted protein was used. Transfection—of this construct produced background binding of Frzbl-HA to the extracellular matrix, both uniform and punctate. Cotransfection of Wnt1CD8 with pcDNA-LacZ showed that transfected cells stained positively for Frzb1-HA and Lacz. Since Wnt1CD8 contains the entire CD8 molecule, a CD8 cDNA was used as an additional negative control. After transfection with Lacz and full-length CE8, Frzbl-HA failed to bind to the transfected cells. Although most of our experiments were carried out at 37°C, Frzbl-HA-conditioned medium also stained WntlCD8-transfected cells after incubation at 4°C for 2 hours.

Attempts to biochemically quantitate the binding of Frzb-1 to Wnt1CD8-transfected cells were unsuccessful due to high\_background binding to control cultures, presumably due to binding to the extracellular matrix. Thus, we were unable to estimate a  $K_D$  for the affinity of the Frzb-1/Wnt-1 interaction. However, when serial dilutions of conditioned medium containing Frzb1-HA were performed (ranging from 2.5 x  $10^{-7}$  to 1.25 x  $10^{-10}$  M), staining of Wnt1CD8-transfected cells was found at all concentrations.

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Although we have been unable to provide biochemical evidence for direct binding between Whats and frzb-1, this cell biological assay indicates that Frzbl-HA can bind, directly or indirectly, to What-1 on the cell membrane in the 10-10 M range.

It is to be understood that while the invention has been described above in conjunction with preferred specific embodiments, the description and examples are intended to illustrate and not limit the scope of the invention, which is defined by the scope of the appended claims.

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#### It is Claimed:

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- 1. A substantially pure protein characterized by a physiologically active form and comprising an amino acid sequence encoded by the DNA of SEQ ID NO:2.
- 2. The protein as in claim 1 having neurotrophic, growth or differentiation factor activity.
- 3. A composition comprising the protein of claim 1 and a physiologically acceptable carrier with which the peptide is admixed.
- 4. An oligonucleotide construct comprising a sequence coding for a protein and an expression vector operatively linked therewith, the protein having neurotrophic, growth or differentiation factor activity and being expressible from SEQ ID NO:2.
- 5. The construct as in claim 4 wherein the expression vector is a mammalian or viral expression vector.
- 6. A substantially pure protein characterized by a physiologically active form and comprising an amino acid sequence encoded by the DNA of SEQ ID NO:4, SEQ ID NO:8, or SEQ ID NO:10.
- 7. The protein as in claim 6 having neurotrophic, growth or differentiation factor activity.
- 8. A composition comprising the protein of claim 6 and a physiologically acceptable carrier with which the protein is admixed.

9. An oligonucleotide construct comprising a sequence coding for a protein and an expression vector operatively linked therewith, the protein being expressible from SEQ ID NO:4, SEQ ID NO:8 or SEQ ID NO:10.

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- 10. The construct as in claim 9 wherein the protein is expressible in soluble form.
- 11. The construct as in claim 9 wherein the expression vector is a mammalian or viral expression vector.
- 12. A complex comprising a substantially pure frzb-1 protein complexed with at least one Wnt protein.
- 13. A substantially pure protein characterized by a physiologically active form and comprising an amino acid sequence encoded by the DNA of SEQ ID NO:6.
- 14. The protein as in claim 13 having mesoderm differentiation activity.
- 15. A composition comprising the protein of claim 13 and a physiologically acceptable carrier with which the protein is admixed.

MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HGNKSARRKA	YNGSRRNIFS	120
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	APQ <u>NTS</u> HGSK	160
AQEIMKEACK	TLPFTQNIVH	ENCDRMVIQN	NLCFGKCISL	200
HVPNQQDRRN	TCSHCLPSKF	TLNHLTLNCT	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1

**SUBSTITUTE** SHEET (RULE 26)

		·				
				ATATCATACA TATAGTATGT		60
	•					
				TGGAGCAGGA		120
TACATGAGTC	CTAGACATAA	TAGCAGACGG	AACACTTACT	ACCTCGTCCT	TTTGTGAGTC	
				AGGTTACTTC		180
TTCCTGCTCT	TTCCTGTTTT	TGTATAAGTG	AATTGTCGTC	TCCAATGAAG	TCTTTTCTTT	
				AGGTCTTGAT		240
CICTOGIGO	ATCCTCGTTC	TAAGACGACC	ACTTATGATT	TCCAGAACTA	CTTGGGGTGT	
TTGGGCATGG	TGATTTTCGC	TTAGTAGCTG	AACTATTTGA	TTCCACCAGA	ACACATACAA	300
				AAGGTGGTCT		
ACAGAAAAGA	GCCAGACATG	AACAAAGTCA	<b>ል</b> ርረጥጥጥጥርጥር	AACAGTTGCC	CATCCAARCA	360
				TTGTCAACGG		360
AAAGTGCAAG	AAGAAAAGCT	TACAATGGTT	CTAGAAGGAA	TATTTTTCCT	CGCCGTTCTT	420
				ataaaaagga		
TTGATAAAAG	AAATACAGAG	GTTACTGAAA	AGCCTGGTGC	CAAGATGTTC	TGGAACAATT	480
				GTTCTACAAG		
TTTTGGTTAA	aatgaatgga	GCCCCACAGA	ATACAAGCCA	TGGCAGTAAA	GCACAGGAAA	540
AAAACCAATT	TTACTTACCT	CGGGGTGTCT	TATGTTCGGT	ACCGTCATTT	CGTGTCCTTT	
				TATTGTACAT		600
ATTACTTTCT	TCGAACGTTT	TGGAACAAAA	AGTGAGTCTT	ATAACATGTA	CTTTTGACAC	
				CATCTCTCTC		660
TGTCCTACCA	CTATGTCTTG	TTAGACACGA	AACCATTTAC	GTAGAGAGAG	GTACAAGGTT	
					ACCCTGAACC	720
TAGTCGTTCT	AGCTGCTTTA	TGAACAAGGG	TAACGAACGG	CAGGTTTAAA	TGGGACTTGG	
					ATGGTAGAGG	780
TGGACTGCGA	CTTAACATGA	CCTAGATTCT	TACATCATTT	CCAACAGTAC	TACCATCTCC	
AATGCACGTG	TGAAGCTCAT	AAGAGCAACT	TCCACCAAAC	TGCACAGTTT	AACATGGATA	840
TTACGTGCAC	ACTTCGAGTA	TTCTCGTTGA	AGGTGGTTTG	ACGTGTCAAA	TTGTACCTAT	
CATCTACTAC	CCTGCACCAT	TAAAGGACTG	CCATACAGTA	TGGAAATGCC	CTTTTGTTGG	900
GTAGATGATG	GGACGTGGTA	ATTTCCTGAC	GGTATGTCAT	ACCTTTACGG	GAAAACAACC	
AATATTTGTT	ACATACTATG	CATCTAAAGC	ATTATGTTGC	CTTCTATTTC	ATATAACCAC	960
TTATAAACAA	TGTATGATAC	GTAGATTTCG	TARTACARCG	GAAGATAAAG	TATATTGGTG	
ATGGARTARG	GATTCTATCA	ATTATAATTA	ACBB RECCO	ቀቀቀጥርጥርጥ <i>እ አ</i>	CATGCAAGAT	1020
TACCTTATTC	CTAACATACT	TAATATTAAT	TGTTTACCGT	AAAACACATT	GTACGTTCTA	1020

## Figure 2A

SUBSTITUTE SHEET (RULE 26)

CTCTGTTCCA TCAGTTGCAA					1080
GAGACAAGGT AGTCAACGTT	CTATTTTCCG	TTATAAACAA	ACTGAAAAAA	AGATGTTTTA	
GAATACCCAA ATATATGATA	AGATAATGGG	GTCAAAACTG	TTAAGGGGTA	ATGTAATAAT	1140
CTTATGGGTT TATATACTAT	TCTATTACCC	CAGTTTTGAC	AATTCCCCAT	TACATTATTA	•
AGGACTAAG TTTGCCCAGG	AGCAGTGACC	CATAACAACC	AATCAGCAGG	TATGATTTAC	1200
TCCCTGATTC AAACGGGTCC	TCGTCACTGG	GTATTGTTGG	TTAGTCGTCC	ATACTAAATG	
TGGTCACCTG TTTAAAAGCA					1260
ACCAGTGGAC AAATTTTCGT	TTGTAGAATA	ACCAACGATA	CCCAATGACG	AAGACCCGTT	
AATGTGTGCC TCATAGGGGG	GTTAGTGTGT	TGTGTACTGA	ATAAATTGTA	TTTATTTCAT	1320
TTACACACGG AGTATCCCCC	CAATCACACA	ACACATGACT	TATTTAACAT	Aaataaagta	
TGTTACAAAA AAAAAAAA					
ACAATGTTTT TTTTTTT	•				

Figure 2B

MSRTRKVDSL	LLLAIPGLAL	LLLPNAYCAS	CEPVRIPMCK	Smpwnmtkmp	NHLHHSTQAN	60
AILAIEQFEG	LLTTECSQDL	LFFLCAMYAP	ICTIDFQHEP	IKPCKSVCER	ARAGCEPILI	120
KYRHTWPESL	ACEELPVYDR	GVCISPEAIV	TVEQGTDSMP	DESMOSNINGN	CGSGREHCKC	180
KPMKATQKTY	LKNNYNYVIR	AKVKEVKVKC	HDATAIVEVK	EILKSSLVNI	PKDTVTLYTN	240
SGCLCPQLVA	NEEYIIMGYE	DKERTRLLLV	EGSLAEKWRD	RLAKKVKRWD	QKLRRPRKSK	300
DPVAPIPNKN	SNSRQARS					

## Figure 3

				TTAGCCCTAT		60
CTTAAGGGAA	AGTGTGTCCT	GAGGACCGTC	TCCACTTACC	AATCGGGATA	CCTAAACCAA	•
					GGATTTTTAT	120
ACAACTAAAA	CTGTGTACTA	ACTAACGAAA	GTCTATCCTA	ACTTCCTGAA	CCTARARATA	•
 CTAATTCTGC	ACTTTTAAAT	TATCTGAGTA	ATTGTTCATT	TTGTATTGGA	TGGGACTAAA	180
				AACATAACCT		
GATAAACTTA	ACTCCTTGCT	TTTGACTTGC	CCATAAACTA	TAAGGTGGGG	TGACTTCTAC	240
				ATTCCACCC		240
<b>ምምር/ምምም</b> ምአ/	<b>み</b> ずごでごごごろ	\$ ##### <b>^</b>	#1 <b>P</b> #0000	ATTCCCTCTA		
				TAAGGGAGAT		300
MCGMARIG	TACACGGGTC	TANANGGGAC	ATAAGGGACA	TAAGGGAGAT	TTCATTCGGA	
ACACATACAG	GTTGGGCAGA	ATAACAATGT	CTCGAACAAG	GAAAGTGGAC	TCATTACTGC	360
TGTGTATGTC	CAACCCGTCT	TATTGTTACA	GAGCTTGTTC	CTTTCACCTG	AGTAATGACG	
TACTGGCCAT	ACCTGGACTG	CCCCTTTCTCT	#18###18### N	TGCTTACTGT	COMMOCMOMO	400
ATGACCGGTA	TGGACCTGAC	CGCGAAGAGA	ATANTGGGTT	ACGAATGACA	CCAACCACAC	420
				CATGACCAAG		480
TCGGACACGC	CTAGGGGTAC	ACGTTTAGAT	ACGGTACCTT	GTACTGGTTC	TACGGGTTGG	
ATCTCCACCA	CAGCACTCAA	GCCAATGCCA	TCCTGGCAAT	TGAACAGTTT	GAAGGTTTGC	540
				ACTTGTCAAA		
				TGCCATGTAT		600
ACTGGTGACT	TACATCGGTC	CTGGAAAACA	AGAAAGACAC	ACGGTACATA	CGGGGGTAAA	
GTACCATCGA	TTTCCAGCAT	GAACCAATTA	AGCCTTGCAA	GTCCGTGTGC	GAAAGGGCCA	660
				CAGGCACACG		
GGGCCGGCTG	TGAGCCCATT	CTCATAAAGT	ACCGGCACAC	TTGGCCAGAG	AGCCTGGCAT	720
CCCGGCCGAC	ACTCGGGTAA	GAGTATTTCA	TGGCCGTGTG	AACCGGTCTC	TOGGACOGTA	
GTGAAGAGCT	GCCCGTATAT	GACAGAGGAG	TCTGCATCTC	CCCAGAGGCT	ATCGTCACAG	780
CACTTCTCGA	CGGGCATATA	CTGTCTCCTC	AGACGTAGAG	GGGTCTCCGA	TAGCAGTGTC	
BCC1101100		2000000000				
1GGAACAAGG	AACAGATTCA	ATGCCAGACT	TCTCCATGGA	TTCAAACAAT	GGAAATTGCG	840
ACCITATION	IIGICIAAGI	TACGGTCTGA	AGAGGTACCT	AAGTTTGTTA	CCTTTAACGC	
GAAGCGGCAG	GGAGCACTGT	AAATGCAAGC	CCATGAAGGC	AACCCAAAAG	ACGTATCTCA	900
CTTCGCCGTC	CCTCGTGACA	TTTACGTTCG	GGTACTTCCG	TTGGGTTTTC	TGCATAGAGT	
						* * -
				GGTGAAAGTG CCACTTTCAC		960
·		INGICIOGIT	TICACTTICT	CCACTTTCAC	TTACGGTGC	
ACGCAACAGC	AATTGTGGAA	GTAAAGGAGA	TTCTCAAGTC	TTCCCTAGTG	AACATTOCTA	1020
					TTGTAAGGAT	

### Figure 4A

SUBSTITUTE SHEET (RULE 26)

				CCCCCAGCTT		1080
TTCTGTGTCA	CTGTGACATG	TGGTTGAGTC	CGACGAACAC	GGGGGTCGAA	CAACGGTTAC	
3CC33T3C3T	አ አ ተሞ አ ጥር ርርርር	TATCABCACA	*******	CAGGCTTCTA	Стастеста	1140
	*			•		1140
TCCTTATGTA	TTARTACCCG	ATACTICIGT	TTCTCGCATG	GTCCGAAGAT	GATCACCTTC	
GATCCTTGGC	CGAAAAATGG	AGAGATCGTC	TTGCTAAGAA	AGTCAAGCGC	TGGGATCAAA	1200
CTAGGAACCG	GCTTTTTACC	TCTCTAGCAG	AACGATTCTT	TCAGTTCGCG	ACCCTAGTTT	
AGCTTCGACG	TCCCAGGAAA	AGCAAAGACC	CCGTGGCTCC	AATTCCCAAC	AAAAACAGCA	1260
TCGAAGCTGC	AGGGTCCTTT	TCGTTTCTGG	GGCACCGAGG	TTAAGGGTTG	TTTTTGTCGT	
ATTCCAGACA	AGCGCGTAGT	TAGACTAACG	GAAAGGTGTA	TGGAAACTCT	ATGGACTTTG	1320
TAAGGTCTGT	TCGCGCATCA	ATCTGATTGC	CTTTCCACAT	ACCTTTGAGA	TACCTGAAAC	
				GCACTACAGC		1380
TTTGATTCTA	AACGTAACAA	CCTTCTCGTT	TTTTCTTTAA	CGTGATGTCG	TGCAATATAA	
CTATTGTTTA	CTACAAGAAG	CTGGTTTAGT	TGATTGTAGT	TCTCCTTTCC	TTCTTTTTT	1440
GATAACAAAT	GATGTTCTTC	GACCAAATCA	ACTAACATCA	AGAGGAAAGG	AAGAAAAAA	
					•	
TTATAACTAT	ATTTGCACGT	GTTCCCAGGC	AATTGTTTTA	TTCAACTTCC	AGTGACAGAG	1500
				AAGTTGAAGG		-
					,	
CAGTGACTGA	ATGTCTCAGC	CTAAAGAAGC	TCAATTCATT	TCTGATCAAC	TAATGGTGAC	1560
				AGACTAGTTG		
AAGTGTTTGA	TACTTGGGGA	AAGTGAACTA	ATTGCAATGG	TAAATCAGAG	AAAAGTTGAC	1620
				ATTTAGTCTC		
				••••		
CAATGTTGCT	TTTCCTGTAG	ATGAACAAGT	GAGAGATCAC	ATTTAAATGA	TGATCACTTT	1680
				TAAATTTACT		
			01010111010			
CCATTTAATA	CTTTCAGCAG	TTTTAGTTAG	ATGACATGTA	GGATGCACCT	AAATCTAAAT	1740
				CCTACGTGGA		_
• • • • • • • • • • • • • • • • • • • •						
ATTTTATCAT	AAATGAAGAG	CTGGTTTAGA	CTGTATGGTC	ACTGTTGGGA	AGGTAAATGC	1800
		<del>-</del>		TGACAACCCT		<del>-</del>
CTACTTTGTC	AATTCTGTTT	TAAAAATTGC	CTAAATAAAT	ATTAAGTCCT	AAATAAAA	1860
GATGAAACAG	TTAAGACAAA	ATTTTTAACG	GATTTATTTA	TAATTCAGGA	TTTATTTTT	
	-				, <del></del>	
AAAAAAAA	AAAAA	•				
TTTTTTTTT	TTTTT					

Figure 4B

SUBSTITUTE SHEET (RULE 26)

MLL	LFRAIPM	LLIGLMVLQT	DCEIAQYYID	EEEPPGTVIA	VLSQHSIFNT	TDIPATNFRL	60
MKQ	FNNSLIG	VRESDGQLSI	MERIDREQIC	RQSLHCNLAL	DVVSFSKGHF	KLLNVKVEVR	120
DIN	DHSPHFP	SEIMHVEVSE-	-SSSVGTRIPL	EIAIDEDVGS	NSIQNFQISN	NSHFSIDVLT	180
RAD	GVKYADL	VLMRELDREI	<b>QPTYIMELLA</b>	MDGGVPSLSG	TAVVNIRVLD	FNDNSPVFER	240
STI	AVDLVED	APLGYLLLEL	HATDDDEGVN	GEIVYGFSTL	ASQEVRQLFK	INSRTGSVTL	300
EGQ	VDFETKQ	TYEFEVQAQD	LGPNPLTATC	KVTVHILDVN	DNTPAITITP	LTTVNAGVAY	360
IPE	TATKENF	IALISTTDRA	SGSNGQVRCT	LYGHEHFKLQ	QAYEDSYMIV	TTSTLDRENI	420
AAY	SLTVVAE	DLGFPSLKTK	KYYTVKVSDE	NDNAPVFSKP	QYEASILENN	APGSYITTVI	480
ard	SDSDQNG	KVNYRLVDAK	VMGQSLTTFV	SLDADSGVLR	AVRSLDYEKL	KQLDFEIEAA	540
DNG	IPQLSTR	VQLNLRIVDQ	NDNCPVITNP	LLNNGSGEVL	LPISAPQNYL	VFQLKAEDSD	600
EGH	nsqlfyt	ILROPSRLFA	INKESGEVFL	KKQLNSDHSE	DLSIVVAVYD	LGRPSLSTNA	660
TVK	FILTDSF	PSNVEVVILQ	PSAEEQHQID	MSIIFIAVLA	GGCALLLLAI	FFVACTCKKK	720
AGE	FKQVPEQ	HGTCNEERLL	STPSPQSVSS	SLSQSESCQL	SINTESENCS	VSSNQEQEQQ	780
TGI	KHSISVP	SYHTSGWHLD	NCAMSISGHS	HMGHISTKVQ	WAKEIVTSMT	VTLILVENOK	840
RRA	LSSQCRH	KPVLNTQMNQ	QGSDMPITIS	ATESTRVQKM	GTAHCNMKRA	IDCLTL	

Figure 5
SUBSTITUTE SHEET (HULE 26)

~	1000000						
G	AATTCCCAG	AGATGAACTC	CTTGAGATTG	TTTTAAATGA	CTGCAGGTCT	GGAAGGATTC	60
C	TAAGGGTC	TCTACTTGAG	GAACTCTAAC	AAAATTTACT	GACGTCCAGA	CCTTCCTAAG	
AC	CATTGCCAC	ACTGTTTCTA	GGCATGAAAA	AACTGCAAGT	TTCAACTTTG	TTTTTGGTGC	120
T	STAACGGTG	TGACAAAGAT	CCGTACTTTT	TTGACGTTCA	AAGTTGAAAC	AAAAACCACG	120
-A:	CTTTGATT	CTTCAAGATG	CTGCTTCTCT	TCAGAGCCAT	TCCAATGCTG	CTGTTGGGAC	180
T	<b>IGAAACTAA</b>	GAAGTTCTAC	GACGAAGAGA	AGTCTCGGTA	AGGTTACGAC	GACAACCCTG	200
TO	SATGGTTTT	ACAAACAGAC	TGTGAAATTG	CCCAGTACTA	CATAGATGAA	GAAGAACCCC	240
AC	CTACCAAAA	TGTTTGTCTG	ACACTTTAAC	GGGTCATGAT	GTATCTACTT	CTTCTTGGGG	
<u></u>		3.5 MMC 03.0 MC	mmcmon or				
C1	CCCCCCCCC	MATIGCAGIG	TIGICACAAC	ACTCCATATT	TAACACTACA	GATATACCTG	300
Gr	ICCG1GACA	TIMACGICAC	AACAGIGITG	TGAGGTATAA	ATTGTGATGT	CTATATGGAC	
CZ	ACCAATTT	CCGTCTAATG	AAGCAATTTA	<b>ል ጥል ል ጥጥረጥር</b> ጥ	TATCGGAGTC	CCTCACACTC	200
G1	TGGTTAAA	GGCAGATTAC	TTCCTTABAT	TATTA ACCCA	ATAGCCTCAG	COLORORORO	360
-			TICGIIMMI	INIIMAGGGA	ATAGCCTCAG	GCACTCTCAC	
A	GGGCAGCT	GAGCATCATG	GAGAGGATTG	ACCGGGAGCA	AATCTGCAGG	CAGTCCCTTC	420
T	CCCGTCGA	CTCGTAGTAC	CTCTCCTAAC	TGGCCCTCGT	TTAGACGTCC	GTCAGGGAAG	120
AC	CTGCAACCT	GGCTTTGGAT	GTGGTCAGCT	TTTCCAAAGG	ACACTTCAAG	CTTCTGAACG	480
T	eacgttgga	CCGAAACCTA	CACCAGTCGA	AAAGGTTTCC	TGTGAAGTTC	GAAGACTTGC	
		•		• *			
T	Gaaagtgga	GGTGAGAGAC	ATTAATGACC	ATAGCCCTCA	CTTTCCCAGT	GAAATAATGC	540
AC	CTTTCACCT	CCACTCTCTG	TAATTACTGG	TATCGGGAGT	GAAAGGGTCA	CTTTATTACG	
A	TGTGGAGGT	GTCTGAAAGT	TCCTCTGTGG	GCACCAGGAT	TCCTTTAGAA	ATTGCAATAG	600
T	CACCTCCA	CAGACTTTCA	AGGAGACACC	CGTGGTCCTA	AGGAAATCTT	TAACGTTATC	
n. 9	~~ x ~ x ~ ~ ~	#CCC#CC110					
# T	COMMONIGI	1GGGTCCAAC	TCCATCCAGA	ACTITCAGAT	CTCAAATAAT	AGCCACTTCA	660
12	CLICIACA	ACCCAGGITG	AGGTAGGTCT	TGAAAGTCTA	GAGTTTATTA	TCGGTGAAGT	
Ċ	こみでかるみでんで	CCTABCCACA	CCACAMOCCO	<b>50111500</b>	101 5551 650		
~	WIIGHTOI	CONTROCTOR	CCAGATGGGG	TGAAATATGC	AGATTTAGTC	TTAATGAGAG	720
`	INNUINCE	CANTIGGICT	CGICIACCC	ACTITATACG	TCTAAATCAG	AATTACTCTC	
Ā	CTGGACAG	GGAAATCCAG	CCARCATACA	TAATCCACCT	ACTAGCAATG	 CNECCCOMO	700
T	GACCTGTC	CCTTTAGGTC	GGTTGTATGT	ATTACCTCA	TGATCGTTAC	CTACCCCALC	780
					IGNICOTING	CINCCUCAC	
T	CCATCACT	ATCTGGTACT	GCAGTGGTTA	ACATCCGAGT	CCTGGACTTT	AATGATAACA	840
A:	rggtagtga	TAGACCATGA	CGTCACCAAT	TGTAGGCTCA	GGACCTGAAA	TTACTATTCT	040
					•		
G	CCCAGTGTT	TGAGAGAAGC	ACCATTGCTG	TGGACCTAGT	AGAGGATGCT	CCTCTGGGAT	900
C	<b>GGTCACAA</b>	ACTCTCTTCG	TGGTAACGAC	ACCTGGATCA	TCTCCTACGA	GGAGACCCTA	-
A	CCTTTTGTT	GGAGTTACAT	GCTACTGACG	ATGATGAAGG	AGTGAATGGA	GAAATTGTTT	960
T	<b>Gaaaaacaa</b>	CCTCAATGTA	CGATGACTGC	TACTACTTCC	TCACTTACCT	CTTTAACAAA	
	•						
A:	rggattcag	CACTTTGGCA	TCTCAAGAGG	TACGTCAGCT	ATTTAAAATT	AACTCCAGAA	1020
T	ACCTAAGTC	GTGAAACCGT	AGAGTTCTCC	ATCCACTCCA	**********	TTC LOCATION	

# Figure 6A SUBSTITUTE SHEET (RULE 26)

CTGGCAGTGT TACTCT GACCGTCACA ATGAGA					1080
AGGTACAAGC CCAAGA TCCATGTTCG GGTTCT					1140
ATATACTTGA TGTAAA TATATGAACT ACATTT					1200
ATGCAGGAGT TGCCTA TACGTCCTCA ACGGAT					1260
GCACTACTGA CAGAGO CGTGATGACT GTCTCG					1320
AGCACTITAA ACTACA TCGTGAAATT TGATGT					1380
TAGACAGGGA AAACAT ATCTGTCCCT TTTGTA					1440
CCTCATTGAA GACCAA GGAGTAACTT CTGGTT					1500
CTGTATTTC TAAACC					1560
ATATAACTAC AGTGAT TATATTGATG TCACTA					1620
GACTTGTGGA TGCAAA CTGAACACCT ACGTTT					1680
ACTCTGGAGT ATTGAG TGAGACCTCA: TAACTC					1740
TTGAAATTGA AGCTGC					1800
TCAGAATAGT TGATCA AGTCTTATCA ACTAG					1860
GCTCGGGTGA AGTTCT CGAGCCCACT TCAAGI					1920
AAGCCGAGGA TTCAGI TTCGGCTCCT AAGTC	ntgaa gggcacaact Pactt cccgtgttga	CCCAGCTGTT GGGTCGACAA	CTATACCATA GATATGGTAT	CTGAGAGATC GACTCTCTAG	1980
CAAGCAGATT GTTTGG GTTCGTCTAA CAAAC	XATT AACAAAGAAA GTAA TTGTTTCTTT	GTGGTGAAGT CACCACTTCA	GTTCCTGAAA CAAGGACTTT	AAACAATTAA TTTGTTAATT	2040
ACTCTGACCA TTCAG TGAGACTGGT AAGTC	AGGAC TTGAGCATAG ICCTG AACTCGTATO	TAGTTGCAGT ATCAACGTCA	GTATGACTTG CATACTGAAC	GGAAGACCTT CCTTCTGGAA	2100
CATTATOCAC CAATG	CTACA GTTAAATTCA GATGT CAATTTAAGT	TCCTCACCGA AGGAGTGGCT	CTCTTTTCCT GAGAAAAGGA	TCTAACGTTG AGATTGCAAC	2160

# Figure 6B SUBSTITUTE SHEET (RULE 26)

AAGTCGTTAT TTTGCAACCA TCTGCAGAAG AGCAGCACCA GATCGATATG TCCATTAT	
TTCAGCAATA AAACGTTGGT AGACGTCTTC TOGTCGTGGT CTAGCTATAC AGGTAATA	ra
TCATTGCAGT GCTGGCTGGT GGTTGTGCTT TGCTACTTTT GGCCATCTTT TTTGTGGC	
AGTAACGTCA CGACCGACCA CCAACACGAA ACGATGAAAA CCGGTAGAAA AAACACCG	3A
GTACTTGTAA AAAGAAAGCT GGTGAATTTA AGCAGGTACC TGAACAACAC GGAACATG	
CATGAACATT TTTCTTTCGA CCACTTAAAT TCGTCCATGG ACTTGTTGTG CCTTGTAC	3T
epoches of the second of the s	
ATGAAGAACG CCTGTTAAGC ACCCCATCTC CCCAGTOGGT CTCTTCTTCT TTGTCTCAC	
TACTTCTTGC GGACAATTCG TGGGGTAGAG GGGTCAGCCA GAGAAGAAGA AACAGAGT	CA C
CTGAGTCATG CCAACTCTCC ATCAATACTG AATCTGAGAA TTGCAGCGTG TCCTCTAAC	C 2460
GACTCAGTAC GGTTGAGAGG TAGTTATGAC TTAGACTCTT AACGTCGCAC AGGAGATTO	GG .
AAGAGCAGCA TCAGCAAACA GGCATAAAGC ACTCCATCTC TGTACCATCT TATCACACI	AT 2520
TTCTCGTCGT AGTCGTTTGT CCGTATTTCG TGAGGTAGAG ACATGGTAGA ATAGTGTG	
CTGGTTGGCA CCTGGACAAT TGTGCAATGA GCATAAGTGG ACATTCTCAC ATGGGGCA	ZA 2580
GACCAACCGT GGACCTGTTA ACACGTTACT CGTATTCACC TGTAAGAGTG TACCCCGT	
TTAGTACAAA GGTACAGTGG GCAAAGGAGA TAGTGACTTC AATGACAGTG ACTCTGATA	AC 2640
AATCATGTTT CCATGTCACC CGTTTCCTCT ATCACTGAAG TTACTGTCAC TGAGACTA	
TAGTGGAGAA TCAGAAAAGA AGAGCATTGA GCAGCCAATG CAGGCACAAG CCAGTGCTG	CA 2700
ATCACCTCTT AGTCTTTTCT TCTCGTAACT CGTCGGTTAC GTCCGTGTTC GGTCACGA	
withouter serential telegistic calculative difficulties calculation	31
ATACACAGAT GAATCAGCAG GGTTCCGACA TGCCGATAAC TATTTCAGCC ACCGAATC	AA 2760
TATGTGTCTA CTTAGTCGTC CCAAGGCTGT ACGGCTATTG ATAAAGTCGG TGGCTTAG	
INTOIGETCIA CITAGLOSIC COAAGGCIGI ACGGCIATIG ATARAGICGG IGGCITAG	r T
CAAGGGTCCA GAAAATGGGA ACTGCACATT GCAATATGAA AAGGGCTATA GACTGTCT	ra 2820
GTTCCCAGGT CTTTTACCCT TGACGTGTAA CGTTATACTT TTCCCGATAT CTGACAGA	
GITCOCAGGI CITITACCCI IGACGIGIAA CGITATACTI TICCCGATAT CIGACAGA	R.T
CTCTGTAGCT CCTGTATATT ACAATACCTA CCATGCAAGA ATGCCTAACC TGCACATA	~ 2004
GAGACATCGA GGACATATAA TGTTATGGAT GGTACGTTCT TACGGATTGG ACGTGTAT	
GAGACATOGA GGACATATAA TGTTATGGAT GGTAGGTTCT TACGGATTGG ACGTGTAT	فاف
GAACCATACC CTTAGAGACC CTTATTACCA TATCAATAAT CCTGTTGCTA ATCGGATG	
CTTGGTATGG GAATCTCTGG GAATAATGGT ATAGTTATTA GGACAACGAT TAGCCTAC	GT
GGCGGAATAT GAAAGAGATT TAGTCAACAG AAGTGCAACG TTATCTCCGC AGAGATCG	
CCGCCTTATA CTTTCTCTAA ATCAGTTGTC TTCACGTTGC AATAGAGGCG TCTCTAGC	<b>A</b> G
TAGCAGATAC CAAGAATTCA ATTACAGTCC GCAGATATCA AGACAGCTTC ATCCTTCA	
ATCCTCTATG GTTCTTAAGT TAATGTCAGG CGTCTATAGT TCTGTCGAAG TAGGAAGT	CT
The state of the s	
AATTGCTACA ACCITTTAAT CATTAGGCAT GCAAGTGAGA ATGCACAAAG GCAAGTGC	TT 3120
TTAACGATGT TGGAAAATTA GTAATCCGTA CGTTCACTCT TACGTGTTTC CGTTCACG	AA
TAGCATGAAA GCTAAATATA TGGAGTCTCC CCTTTCCCTC TGATGGATGG GGGGAGAC	
ATCGTACTTT CGATTTATAT ACCTCAGAGG GGAAAGGGAG ACTACCTACC CCCCTCTG	TG
AGGACAGTGC ATAAATATAC AGCTGCTTTC TATTTGCATT TCACTTGGGA ATTTTTTG	
TCCTGTCACG TATTTATATG TCGACGAAAG ATAAACGTAA AGTGAACCCT TAAAAAAC	AA
TTTTTTACAT ATTTATTTTT CCTGAATTGA ATGTGACATT GTCCTGTCAC CTAACTAG	
AAAAAATGTA TAAATAAAAA GGACTTAACT TACACTGTAA CAGGACAGTG GATTGATC	GT
interest the transfer of the t	

## Figure 6C

SUBSTITUTE SHEET (RULE 26)

### 11/18

ATTARATOCA CAGACCTACA GTCARATATT TGAGGGCCCC TGARACAGCA CATCAGTCAG TARTTTAGGT GTCTGGATGT CAGTTTATAR ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC	3360
GACCTARAGT GGCCTTTTTA CTTTTAGCAG CTCCTGGGTC TGCCCTCTGT GTTAATCAGC CTGGATTTCA CCGGARARAT GARARTCGTC GAGGACCCAG ACGGGAGACA CARTTAGTCG	3420
CCCTGGTCAA GTCCTGAGTA GGATCATGGC GTTTTTATAT GCATCTCACC TACTTTGGAC GGGACCAGTT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG	3480
GTGATTTACA CATAATAGGA AACGCTTGGT TTCAGTGAAG TCTGTGTTGT ATATATCTG CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAACA TATATAAGAC	3540
TTATATACAC GCATTTTGTG TTTGTGTATA TATTTCAAGT CCATTCAGAT ATGTGTATAT AATATATGTG CGTAAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA	3600
AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTTCC TCAATAAATA TTTAAAT TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAAGG AGTTATTTAT AAATTTA	

SUBSTITUTE SHEET (RULE 26)

MVCCGPGRML	LGWAGLLVLA	ALCLLQVPGA	QAAACEPVRI	PLCKSLPWNM	TKMPNHLHHS	60
TQANAILAME	QFEGLLGTHC	SPDLLFFLCA	MYAPICTIDF	QHEPIKPCKS	VCERARQGCE	120
PILIKYRHSW	PESLACDELP	VYDRGVCISP	EAIVTADGAD	FPMDSSTGHC	RGASSERCKC	180
KPVRATQKTY	FRNNYNYVIR	AKVKEVKMKC	HDVTAVVEVK	EILKASLVNI	PRDTVNLYTT	240
SGCLCPPLTV	NEEYVIMGYE	DEERSRLLLV	EGSIAEKWKD	RLGKKVKRWD	MKLRHLGLGK	300
TDASDSTQNQ	KSGRNSNPRP	ARS.				

AAGCCTGGGA	CCATGGTCTG	CTGCGGCCCG	GGACGGATGC	TGCTAGGATG	GGCCGGGTTG	60
TTCGGACCCT	GGTACCAGAC	GACGCCGGGC	CCTGCCTACG	ACGATCCTAC	CCGGCCCAAC	
					,	
СТАСТССТСС	CTGCTCTCTG	ССТССТССАС	СТСССССАС	СТСАСССТСС	AGCCTGTGAG	120
	GACGAGAGAC					120
ONICHOUNCE	GACGAGAGAC	GGACGAGGIC	CACGGGCCIC	GAGICCGACG	ICOGNONCIC	•
OCTOTOTO CO CO	maaaaaamama	CA A CONCOCODO	COCTOCALACIA	maxaaxaa	0000110010	100
	TCCCGCTGTG					180
GGACAGGCGT	AGGGCGACAC	G1 TCAGGGAA	GGGACCTTGT	ACTGGTTCTA	CGGGTTGGTG	
					-	
	GCACCCAGGC	· · · · · · · · · · · · ·				240
GACGTGGTGT	CGTGGGTCCG	ATTGCGGTAG	GACCGGTACC	TTGTCAAGCT	TCCCGACGAC	
					*	
GGCACCCACT	GCAGCCCGGA	TCTTCTCTTC	TTCCTCTGTG	CAATGTACGC	ACCCATTTGC	300
CCGTGGGTGA	CGTCGGGCCT	AGAAGAGAAG	AAGGAGACAC	GTTACATGCG	TGGGTAAACG	
ACCATCGACT	TCCAGCACGA	GCCCATCAAG	CCCTGCAAGT	CTGTGTGTGA	GCGCGCCCGA	360
	AGGTCGTGCT					
100111001011		00001110110	000110011011	0.101.01.01.01	000000001	
CAGGGGTGCG	AGCCCATTCT	CATCAACTAC	CCCC3 CTCCT	CCCCCCAAAC	COMPAGE	420
·	TCGGGTAAGA					420
GICCCGACGC	ICGGGIAAGA	GIAGIICAIG	GCGGIGAGCA	CCGGCCTTTC	GAACCGGACG	
	000000000000000000000000000000000000000			cmc> cccc> m	000010000	400
	CGGTGTACGA					480
CIGCICGACG	GCCACATGCT	GGCGCCGCAC	ACGTAGAGAG	GACTCCGGTA	GCAGTGGCGC	
		•				
	ATTITICCTAT					540
CIGCCICGCC	TAAAAGGATA	CCTAAGTTCA	TGACCTGTGA	CGTCTCCCCG	TTCGTCGCTT	,
CGTTGCAAAT	GTAAGCCTGT	CAGAGCTACA	CAGAAGACCT	ATTTCCGGAA	CAATTACAAC	600
GCAACGTTTA	CATTCGGACA	GTCTCGATGT	GTCTTCTGGA	TAAAGGCCTT	GTTAATGTTG	
			•			
TATGTCATCC	GGGCTAAAGT	TAAAGAGGTA	AAGATGAAAT	GTCATGATGT	GACCGCCGTT	660
ATACAGTAGG	CCCGATTTCA	ATTTCTCCAT	TTCTACTTTA	CAGTACTACA	CTGGCGGCAA	
GTGGAAGTGA	AGGAAATTCT	AAAGGCATCA	СТССТАВАСА	TTCCAAGGGA	CACCGTCAAT	720
	TCCTTTAAGA			= ' -		
CACCIICACI	ICCITIANGA	IIICCGIAGI	GACCATITGE	Magriceci	01000101111	
COORDANACOA	CCTCTGGCTG	~~m~m~~~		MONNING CON	3 m3 m~m~3 m~	780
						. 700
GAMATATGGT	GGAGACCGAC	GGAGACAGGA	GGTGAATGAC	AGTIACICCI	ININCAGING	
1 = C C C C = T = T = C		1 00000001 00		m1 01 1 00 000	m1 m1 comc1 c	840
					TATAGCTGAG	840
TACCCGATAC	TTCTGCTCCT	TGCAAGGTCC	AATGAGAACC	ATCTTCCGAG	ATATCGACTC	
					CCGACACCTT	900
TTCACCTTCC	TAGCCGAACC	ATTCTTTCAC	TTCGCGACCC	TATACTTTGA	GGCTGTGGAA	
GGACTGGGT	AAACTGATG(	TAGCGATTC	ACTCAGAATC	AGAAGTCTGG	CAGGAACTCT	960
CCTGACCCAT	TTTGACTAC	ATCGCTAAGO	G TGAGTCTTAG	TCTTCAGACO	GTCCTTGAGA	

Figure 8A SUBSTITUTE SHEET (RULE 26)

AATCCCCGGC CA				1020
TTCTAAGACT GO				1080
TTGTTTACCG CA				1140
CTTAATGGCG TY GAATTACCGC AC				1200
GGGACTGTTC TO				1260
CTGGACTCCC TY GACCTGAGGG A			 	1320
TAAAGAGAGA A'	 		 	1380
GCTGCGCTTA T			 ,	1440
ATACATGTTT A	 		 	1500
CCAACACCAG G GGTTGTGGTC C	 		 	1560
CAGGCAGCAA A GTCCGTCGTT T		•		1620
CACACTGGAA T	 			1680
TTTGTTCATA A AAACAAGTAT T	 		 	1740
ATCTCTATAG C	 			1800

Figure 8B
SUBSTITUTE SHEET (RULE 26)

## 15/18

	TTGGCTTGCT AACCGAACGA				1860
	GTGTTATTTA CACAATAAAT		 		1920
	GTGCACATTT CACGTGTAAA				1980
	TGTGTTTATG ACACAAATAC			CCATTGCACA GGTAACGTGT	2040
	ACTAGATTAG TGATCTAATC		 		2100
	TAATGCTCCA ATTACGAGGT			TCAACAGAGA AGTTGTCTCT	2160
CGACAACAAC GCTGTTGTTG		·			·.

Figure 8C SUBSTITUTE SHEET (RULE 26)

MVCGSPGGML	LLRAGLLALA	ALCLLRVPGA	RAAACEPVRI	PLCKSLPWNM	TKMPNHLHHS	60
TQANAILAIE	QFEGLLGTHC	SPDLLFFLCA	MYAPICTIDF	QHEPIKPCKS	VCERARQGCE	120
PILIKYRHSW	PENLACEELP	VYDRGVCISP	EAIVTADGAD	FPMDSSNGNC	RGASSERCKC	18
KPIRATOKTY	FRNNYNYVIR	AKVKEIKTKC	HDVTAVVEVK	EILKSSLVNI	PRDTVNLYTS	24
SGCLCPPLNV	NEEYIIMGYE	DEERSRLLLV	EGSIAEKWKD	RLGKKVKRWD	MKLRHLGLSK	30
SDSSNSDSTO	CUKCCBNGND	POARN				

Figure 9 SUBSTITUTE SHEET (RULE 26)

GGCGGAGCGG	GCCTTTTGGC	GTCCACTGCG	CGGCTGCACC	CTGCCCCATC	TGCCGGGATC	60
CCGCCTCGCC	CGGAAAACCG	CAGGTGACGC	GCCGACGTGG	GACGGGGTAG	ACGGCCCTAG	
ATGGTCTGCG	GCAGCCCGGG	AGGGATGCTG	CTGCTGCGGG	CCGGGCTGCT	TGCCCTGGCT	120
TACCAGACGC	CGTCGGGCCC	TCCCTACGAC	GACGACGCCC	GGCCCGACGA	ACGGGACCGA	
GCTCTCTGCC	TGCTCCGGGT	GCCCGGGGCT	CGGGCTGCAG	CCTGTGAGCC	CGTCCGCATC	180
CGAGAGACGG	ACGAGGCCCA	CGGGCCCCGA	GCCCGACGTC	GGACACTCGG	GCAGGCGTAG	
CCCCTGTGCA	AGTCCCTGCC	CTGGAACATG	ACTAAGATGC	CCAACCACCT	GCACCACAGC	240
GGGGACACGT	TCAGGGACGG	GACCTTGTAC	TGATTCTACG	GGTTGGTGGA	CGTGGTGTCG	
	ACGCCATCCT					300
TGAGTCCGGT	TGCGGTAGGA	CCGGTAGCTC	GTCAAGCTTC	CAGACGACCC	GTGGGTGACG	
	TGCTCTTCTT					360
TCGGGGCTAG	ACGAGAAGAA	GGAGACACGG	TACATGCGCG	GGTAGACGTG	GTAACTGAAG	
CAGCACGAGC	CCATCAAGCC	CTGTAAGTCT	GTGTGCGAGC	GGGCCCGGCA	GGGCTGTGAG	420
GTCGTGCTCG	GGTAGTTCGG	GACATTCAGA	CACACGCTCG	CCCGGGCCGT	CCCGACACTC	
	TCAAGTACCG					480
GGGTATGAGT	AGTTCATGGC	GGTGAGCACC	GGCCTCTTGG	ACCGGACGCT	CCTCGACGGT	
GTGTACGACA	GGGCGTGTG	CATCTCTCCC	GAGGCCATCG	TTACTGCGGA	CGGAGCTGAT	540
CACATGCTGT	CCCCGCACAC	GTAGAGAGGG	CTCCGGTAGC	AATGACGCCT	GCCTCGACTA	
	ATTCTAGTAA					600
AAAGGATACC	TAAGATCATT	GCCTTTGACA	TCTCCCCGTT	CGTCACTTGC	GACATTTACA	
	GAGCTACACA					660
TTCGGATAAT	CTCGATGTGT	CTTCTGGATA	AAGGCCTTGT	TAATGTTGAT	ACAGTAAGCC	
GCTAAAGTTA	AAGAGATAAA	GACTAAGTGC	CATGATGTGA	CTGCAGTAGT	GGAGGTGAAG	720
CGATTTCAAT	TTCTCTATTT	CTGATTCACG	GTACTACACT	GACGTCATCA	CCTCCACTTC	
GAGATTCTAA	AGTCCTCTCT	GGTAAACATT	CCACGGGACA	CTGTCAACCT	CTATACCAGC	780
CTCTAAGATT	TCAGGAGAGA	CCATTTGTAA	GGTGCCCTGT	GACAGTTGGA	GATATGGTCG	
TCTGGCTGCC	TCTGCCCTCC	ACTTAATGTT	AATGAGGAAT	ATATCATCAT	GGGCTATGAA	840
AGACCGACGG	AGACGGGAGG	TCAATTACAA	עוזארייריואין עוזאר	ጥልጥልርጥልርጥል	CCCCAMACTO	

Figure 10A SUBSTITUTE SHEET (RULE 26)

GATGAGGAAC GTTCCAGATT ACTCTTGGTG GAAGGCTCTA TAGCTGAGAA GTGGAAGGAT 900 CTACTCCTTG CAAGGTCTAA TGAGAACCAC CTTCCGAGAT ATCGACTCTT CACCTTCCTA CGACTCGGTA AAAAAGTTAA GCGCTGGGAT ATGAAGCTTC GTCATCTTGG ACTCAGTAAA 960 GCTGAGCCAT TTTTTCAATT CGCGACCCTA TACTTCGAAG CAGTAGAACC TGAGTCATTT AGTGATTCTA GCAATAGTGA TTCCACTCAG AGTCAGAAGT CTGGCAGGAA CTCGAACCCC 1020 TCACTAAGAT CGTTATCACT AAGGTGAGTC TCAGTCTTCA GACCGTCCTT GAGCTTGGGG CGGCAAGCAC GCAACTAAAT CCCGAAATAC AAAAAGTAAC ACAGTGGACT TCCTATTAAG 1080 GCCGTTCGTG CGTTGATTTA GGGCTTTATG TTTTTCATTG TGTCACCTGA AGGATAATTC ACTTACTTGC ATTGCTGGAC TAGCAAAGGA AAATTGCACT ATTGCACATC ATATTCTATT 1140 TGAATGAACG TAACGACCTG ATCGTTTCCT TTTAACGTGA TAACGTGTAG TATAAGATAA GTTTACTATA AAAATCATGT GATAACTGAT TATTACTTCT GTTTCTCTTT TGGTTTCTGC 1200 CAAATGATAT TTTTAGTACA CTATTGACTA ATAATGAAGA CAAAGAGAAA ACCAAAGACG TTCTCTCTC TCTCAACCCC TTTGTAATGG TTTGGGGGCA GACTCTTAAG TATATTGTGA 1260 AAGAGAGAAG AGAGTTGGGG AAACATTACC AAACCCCCGT CTGAGAATTC ATATAACACT GTTTTCTATT TCACTAATCA TGAGAAAAAC TGTTCTTTTG CAATAATAAT AAATTAAACA 1320 CAAAAGATAA AGTGATTAGT ACTCTTTTTG ACAAGAAAAC GTTATTATTA TTTAATTTGT TGCTGTTACC AGAGCCTCTT TGCTGAGTCT CCAGATGTTA ATTTACTTTC TGCACCCCAA 1380 ACGACAATGG TCTCGGAGAA ACGACTCAGA GGTCTACAAT TAAATGAAAG ACGTGGGGTT TTGGGAATGC AATATTGGAT GAAAAGAGAG GTTTCTGGTA TTCACAGAAA GCTAGATATG 1440 AACCCTTACG TTATAACCTA CTTTTCTCTC CAAAGACCAT AAGTGTCTTT CGATCTATAC CCTTAAAACA TACTCTGCCG ATCTAATTAC AGCCTTATTT TTGTATGCCT TTTGGGCATT 1500 GGAATTTTGT ATGAGACGGC TAGATTAATG TCGGAATAAA AACATACGGA AAACCCGTAA CTCCTCATGC TTAGAAAGTT CCAAATGTTT ATAAAGGTAA AATGGCAGTT TGAAGTCAAA 1560 GAGGAGTACG AATCTTTCAA GGTTTACAAA TATTTCCATT TTACCGTCAA ACTTCAGTTT TGTCACATAG GCAAAGCAAT CAAGCACCAG GAAGTGTTTA TGAGGAAACA ACACCCAAGA 1620 ACAGTGTATC CGTTTCGTTA GTTCGTGGTC CTTCACAAAT ACTCCTTTGT TGTGGGTTCT TGAATTATTT TTGAGACTGT CAGGAAGTAA AATAAATAGG AGCTTAAGAA AGAACATTTT 1680 ACTTAATAAA AACTCTGACA GTCCTTCATT TTATTTATCC TCGAATTCTT TCTTGTAAAA GCCTGATTGA GAAGCACAAC TGAAACCAGT AGCCGCTGGG GTGTTAATGG TAGCATTCTT 1740 CGGACTAACT CTTCGTGTTG ACTTTGGTCA TCGGCGACCC CACAATTACC ATCGTAAGAA CTTTTGGCAA TACATTGAT TTGTTCATGA ATATATTAAT CAGCATTAGA GAAATGAATT 1800 GAAAACCGTT ATGTAAACTA AACAAGTACT TATATAATTA GTCGTAATCT CTTTACTTAA ATAACTAGAC ATCTGCTGTT ATCACCATAG TTTTGTTTAA TTTGCTTCCT TTTAAATAAA TATTGATCTG TAGACGACAA TAGTGGTATC AAAACAAATT AAACGAAGGA AAATTTATTT CCCATTGGTG AAAGTCAAAA AAAAAAAAA AAA GGGTAACCAC TTTCAGTTTT TTTTTTTTTT TTT

Figure 10B SUBSTITUTE SHEET (RULE 26)

#### INTERNATIONAL SEARCH REPORT

International application No.
PCT/US97/10942

A. CLASSIFICATION OF SUBJECT MATTER  IPC(6): Please See Extra Sheet. US CL: 530/300, 350; 514/2; 536/23.1  According to International Patent Classification (IPC) or to both B. FIELDS SEARCHED  Minimum documentation searched (classification system followe U.S.: 530/300, 350; 514/2; 536/23.1  Documentation searched other than minimum documentation to the  Electronic data base consulted during the international search (no DIALOG (MEDLINE, BIOSIS, EMBASE, WPI, USPATFUE xenopus	d by classification symbols)  e extent that such documents are included  ame of data base and, where practicable,	search terms used)
C. DOCUMENTS CONSIDERED TO BE RELEVANT		
Category* Citation of document, with indication, where a	ppropriate, of the relevant passages	Relevant to claim No.
Y, P  BOUWMEESTER et al. Cerberus is factor expressed in the anterior organizer. Nature. 15 August 19 pages 595-601, see entire docum	endoderm of Spemann's 196, Vol. 382, No. 6592,	1-15
Further documents are listed in the continuation of Box C	C. See patent family annex.	
Special estegories of cited documents:  'A" document defining the general state of the art which is not considered to be of particular relevance  'E" earlier document published on or after the international filing date  'L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)  'O" document referring to an oral disclosure, use, exhibition or other means  'P" document published prior to the international filing date but later than the priority date claimed  Date of the actual completion of the international search  29 AUGUST 1997	"T" later document published after the interdate and not in conflict with the applied principle or theory underlying the inventor of particular relevance; the considered novel or cannot be considered to the document is taken alone.  "Y" document of particular relevance; the considered to involve an inventive combined with one or more other such being obvious to a person skilled in the document member of the same patent.  Date of mailing of the international sea 11 SEP 1997	claimed invention cannot be ed to involve an inventive step to claimed invention cannot be step when the document is a documents, such combination e art
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231 Facsimile No. (703) 305-3230	Authorized officer  HEATHER BAKALYAR  Telephone No. (703) 308-0196	Ste

#### INTERNATIONAL SEARCH REPORT

International application No. PCT/US97/10942

A. CLASSIFICATION OF SUBJECT MATTER: IPC (6):
A01N 37/18; A61K 38/00; C07K 1/00, 2/00, 4/00, 7/00, 14/00, 16/00, 17/00; C07H 21/02, 21/04

Form PCT/ISA/210 (extra sheet)(July 1992)\*